

OIPE ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/074,695

DATE: 06/18/2002 TIME: 09:07:03

Input Set : A:\213542000102 Seq List.txt Output Set: N:\CRF3\06182002\J074695.raw

		SEQUENCE LISTING	
		RAL INFORMATION:	
10		APPLICANT: Olsson, Lennart	
11		Naranda, Tatjana	
13	(ii)	TITLE OF INVENTION: RECEPTOR DERIVED PEPTIDES INVOLVED IN	
14		MODULATION OF RESPONSE TO LIGAND BINDI	NG
16	(iii)	NUMBER OF SEQUENCES: 12	
18	(iv)	CORRESPONDENCE ADDRESS:	
19		(A) ADDRESSEE: Morrison and Foerster LLP	
20		(B) STREET: 755 Page Mill Road	
21		(C) CITY: Palo Alto	
22		(D) STATE: California	
23	٠.	(E) COUNTRY: United States	
24		(F) ZIP: 94304-1018	
26	(V)	COMPUTER READABLE FORM:	
27		(A) MEDIUM TYPE: Floppy disk	
28		(B) COMPUTER: IBM PC compatible	
29		(C) OPERATING SYSTEM: PC-DOS/MS-DOS	
30		(D) SOFTWARE: PatentIn Release #1.0, Version #1.30	
32	(vi)	CURRENT APPLICATION DATA:	
33		(A) APPLICATION NUMBER: US/10/074,695	
34		(B) FILING DATE: 11-Feb-2002	
35		(C) CLASSIFICATION:	
45	(vii)	PRIOR APPLICATION DATA:	
38		(A) APPLICATION NUMBER: US 08/788,820	
39	•	(B) FILING DATE: 23-JAN-1997	
42		(A) APPLICATION NUMBER: US 08/701,382	
43		(B) FILING DATE: 22-AUG-1996	
46		(A) APPLICATION NUMBER: US 08/612,999	
47		(B) FILING DATE: 08-MAR-1996	
49	(viii)	ATTORNEY/AGENT INFORMATION:	
50		(A) NAME: Apple, Randolph T.	
51		(B) REGISTRATION NUMBER: 36,429	
52		(C) REFERENCE/DOCKET NUMBER: 213542000102	
54	(ix)	TELECOMMUNICATION INFORMATION:	
55		(A) TELEPHONE: (650) 813-5933	
56		(B) TELEFAX: (415) 494-0792	
57		(C) TELEX;	
60	(2) INFOR	MATION FOR SEQ ID NO: 1:	
62		SEQUENCE CHARACTERISTICS:	
63		(A) LENGTH: 25 amino acids	
64		(B) TYPE: amino acid	

(C) STRANDEDNESS: unknown

65

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```
(D) TOPOLOGY: unknown
66
68
       (ii) MOLECULE TYPE: protein
       (ix) FEATURE:
70
71
             (A) NAME/KEY: CDS
             (B) LOCATION: 50..1030
72
74
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1
       Glu Arg Glu Thr Gln Ile Ala Lys Gly Asn Glu Gln Ser Phe Arg Val
76
77
                       5
                                            10
79
      Asp Leu Arg Thr Leu Leu Arg Tyr Tyr
80
                   20
                                        25
83 (2) INFORMATION FOR SEQ ID NO: 2:
       (i) SEQUENCE CHARACTERISTICS:
85
             (A) LENGTH: 23 amino acids
86
             (B) TYPE: amino acid
87
88
             (C) STRANDEDNESS: unknown
             (D) TOPOLOGY: unknown
89
91
       (ii) MOLECULE TYPE: protein
93
       (ix) FEATURE:
             (A) NAME/KEY: CDS
94
             (B) LOCATION: 50..1030
95
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2
97
       Thr Trp Leu Gly Arg Gln Gly Pro Glu Gly Pro Ser Ser Ile Pro Pro
99
100
                        5
                                             10
102
       Gly Thr Leu Thr Thr Leu Trp
103
                    20
106 (2) INFORMATION FOR SEO ID NO: 3:
108
         (i) SEQUENCE CHARACTERISTICS:
109
              (A) LENGTH: 24 amino acids
110
              (B) TYPE: amino acid
              (C) STRANDEDNESS: unknown
111
112
              (D) TOPOLOGY: unknown
114
        (ii) MOLECULE TYPE: protein
116
        (ix) FEATURE:
117
              (A) NAME/KEY: CDS
118
              (B) LOCATION: 50..1030
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3
120
122
        Lys Thr Asp Ser Gln Ile Leu Lys Glu Leu Glu Glu Ser Ser Phe Arg
                        5
                                             10
                                                                  15
123
        1
125
        Lys Thr Phe Glu Asp Tyr Leu His
126
                    20
129 (2) INFORMATION FOR SEQ ID NO: 4:
         (i) SEQUENCE CHARACTERISTICS:
131
132
              (A) LENGTH: 25 amino acids
133
              (B) TYPE: amino acid
134
              (C) STRANDEDNESS: unknown
135
              (D) TOPOLOGY: unknown
137
        (ii) MOLECULE TYPE: protein
139
        (ix) FEATURE:
```

(A) NAME/KEY: CDS

140

DATE: 06/18/2002

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                 Input Set : A:\213542000102 Seq List.txt
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141
               (B) LOCATION: 50..1030
1/3
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4
145
        Gly Arg Gly Asn Glu Lys Lys Pro Ser Ser Val Arg Ala Leu Ser Tle
146
                         5
                                              10
148
        Val Leu Pro Ile Val Leu Leu Val Phe
149
                     20
152 (2) INFORMATION FOR SEQ ID NO: 5:
154
          (i) SEQUENCE CHARACTERISTICS:
155
               (A) LENGTH: 24 amino acids
156
               (B) TYPE: amino acid
157
               (C) STRANDEDNESS: unknown
158
               (D) TOPOLOGY: unknown
160
        (ii) MOLECULE TYPE: protein
162
        (ix) FEATURE:
163
               (A) NAME/KEY: CDS
164
               (B) LOCATION: 50..1030
166
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5
168
        Lys Thr Glu Ala Glu Lys Gln Ala Glu Lys Glu Glu Ala Glu Tyr Arg
169
                         5
                                              10
171
        Lys Val Phe Glu Asn Phe Leu His
172
                     20
175 (2) INFORMATION FOR SEQ ID NO: 6:
177
         (i) SEQUENCE CHARACTERISTICS:
178
               (A) LENGTH: 23 amino acids
179
               (B) TYPE: amino acid
180
               (C) STRANDEDNESS: unknown
181
               (D) TOPOLOGY: unknown
183
        (ii) MOLECULE TYPE: protein
185
        (ix) FEATURE:
186
              (A) NAME/KEY: CDS
187
              (B) LOCATION: 50..1030
189
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6
191
        Lys Lys Glu Asn Lys Ile Val Pro Ser Lys Glu Ile Val Trp Trp Met
192
        1
                         5
                                             10
                                                                  15
194
        Asn Leu Ala Glu Lys Ile Pro
195
                    20
198 (2) INFORMATION FOR SEQ ID NO: 7:
200
         (i) SEQUENCE CHARACTERISTICS:
201
              (A) LENGTH: 25 amino acids
202
              (B) TYPE: amino acid
203
              (C) STRANDEDNESS: unknown
204
              (D) TOPOLOGY: unknown
206
        (ii) MOLECULE TYPE: protein
208
        (ix) FEATURE:
209
              (A) NAME/KEY: CDS
210
              (B) LOCATION: 50..1030
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7
212
214
        Glu Lys Lys Pro Val Pro Trp Glu Ser His Asn Ser Ser Glu Thr Cys
215
                                             10
                                                                  15
```

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Input Set : A:\213542000102 Seq List.txt
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```
217
         Gly Leu Pro Thr Leu Val Gln Thr Tvr
218
                     20
                                          25
221 (2) INFORMATION FOR SEQ ID NO: 8:
          (i) SEQUENCE CHARACTERISTICS:
224
               (A) LENGTH: 24 amino acids
225
               (B) TYPE: amino acid
226
               (C) STRANDEDNESS: unknown
227
               (D) TOPOLOGY: unknown
         (ii) MOLECULE TYPE: protein
229
231
         (ix) FEATURE:
232
               (A) NAME/KEY: CDS
233
               (B) LOCATION: 50..1030
         (xi) SEQUENCE DESCRIPTION: SEO ID NO: 8
235
237
        Tyr Lys Glu Gly Thr Met Leu Asn Cys Glu Cys Lys Thr Gly Phe Arq
238
        1
                         5
                                              10
240
        Arg Ile Lys Ser Gly Ser Leu Tyr
241
                     20
244 (2) INFORMATION FOR SEQ ID NO: 9:
246
         (i) SEQUENCE CHARACTERISTICS:
247
               (A) LENGTH: 24 amino acids
248
               (B) TYPE: amino acid
249
               (C) STRANDEDNESS: unknown
250
               (D) TOPOLOGY: unknown
252
        (ii) MOLECULE TYPE: protein
254
        (ix) FEATURE:
255
               (A) NAME/KEY: CDS
256
               (B) LOCATION: 50..1030
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9
258
260
        Leu Leu Glu Gly Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile
261
        1
                        5
                                              10
                                                                   15
263
        Gln Cys His Pro Glu Cys Leu Pro
264
                    20
267 (2) INFORMATION FOR SEQ ID NO: 10:
269
         (i) SEQUENCE CHARACTERISTICS:
270
              (A) LENGTH: 27 amino acids
271
              (B) TYPE: amino acid
272
              (C) STRANDEDNESS: unknown
273
              (D) TOPOLOGY: unknown
275
        (ii) MOLECULE TYPE: protein
277
        (ix) FEATURE:
278
              (A) NAME/KEY: CDS
279
              (B) LOCATION: 50..1030
281
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10
283
        Pro Lys Thr Asp Ser Gln Ile Leu Lys Glu Leu Glu Glu Ser Ser Phe
284
                                             10
286
        Arg Lys Thr Phe Glu Asp Tyr Leu His Asn Val
287
                    20
290 (2) INFORMATION FOR SEQ ID NO: 11:
        (i) SEQUENCE CHARACTERISTICS:
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Input Set : A:\213542000102 Seg List.txt Output Set: N:\CRF3\06182002\J074695.raw

```
293
              (A) LENGTH: 36 amino acids
294
              (B) TYPE: amino acid
295
              (C) STRANDEDNESS: unknown
296
              (D) TOPOLOGY: unknown
298
        (ii) MOLECULE TYPE: protein
300
        (xi) SEQUENCE DESCRIPTION: SEO ID NO: 11
302
        Gly Asn Glu Gln Ser Phe Arg Val Asp Leu Arg Thr Leu Leu Arg Tyr
303
        1
                     5
                                           10
                                                               15
305
        Ala Gly Gly Asn Glu Gln Ser Phe Arg Val Asp Leu Arg Thr Leu
306
                   20
                                        25
                                                            30
308
        Leu Arg Tyr Ala
309
                35
312 (2) INFORMATION FOR SEO ID NO: 12:
314
         (i) SEQUENCE CHARACTERISTICS:
315
              (A) LENGTH: 28 amino acids
316
              (B) TYPE: amino acid
317
              (C) STRANDEDNESS: unknown
318
              (D) TOPOLOGY: unknown
320
        (ii) MOLECULE TYPE: protein
322
        (ix) FEATURE:
323
              (A) NAME/KEY: CDS
324
              (B) LOCATION: 50..1030
326
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12
328
       Glu Tyr Glu Leu Gln Tyr Lys Glu Val Asn Glu Thr Lys Trp Lys Met
329
       1
                   5
                                          10
                                                                15
331
       Met Asp Pro Ile Leu Thr Thr Ser Val Pro Val Tyr
332
                   20
                                        25
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/074,695

DATE: 06/18/2002 TIME: 09:07:04

Input Set : A:\213542000102 Seq List.txt
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L:33 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:34 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]